### PATENT COOPERATION TREATY

## **PCT**

PORT PC7

JUM 1,00

INTERNATIONAL PRELIMINARY EXAMINATION REPORT

(PCT Article 36 and Rule 70)

12

Applicant's or agent's file reference	FOR FURTHER ACTION	See Norit	ication of Transmittal of International
PCT 99-45	FOR FURTHER ACTION	Preliminar	y Examination Report (Form PCT/IPEA/416
International application No.	International filing date (day)	(month/year)	Priority date (day/month/year)
PCT/US00. 19007	13 JULY 2000		13 JULY 1999
	IPC: or national classification and I +35/440	PC	
Applicant MIDWEST RESEARCH INSTIT	UTE		
Examining Authority ar  2. This REPORT consists and a been amended and a	nd is transmitted to the applicant of a total of sheets.  companied by ANNEXES, i.e., she	according to ets of the desc eets containin	ription, claims and or drawings which hav 12 rectifications made before this Authority
These annexes consist of	a total of <u>C</u> sheets.		
3. This report contains indica	itions relating to the following it	ems:	
I X Basis of the	report		
H Priority			
III Non-establish	nment of report with regard to no	velty, invent	ive step or industrial applicability
<b></b> -	v of invention	·	11 332
V X Reasoned state		ard to novelty, ent	inventive step or industrial applicability;
VI Certain docume	ents cited		
VII Certain defects	in the international application		
	ations on the international applicate		
VIII VIIIIII VIIII VIII	don's on the international approach	OII	
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Name and mailing address of the IP	EA US Autho	orized officer	
Commissioner of Patents and Tr. Box PCT		fair	Bridge for
Washington, D.C. 20231		ANJUNATH	RAO
[acsimile No. (703):305-3230	Telep	hone No. 💢 🚓	(03) 308-0196

## INTERNATIONAL PRELIMINARY EXAMINATION REPORT

International application No.

PCT/US00/19007

I. Basis of the report	
1. With regard to the elements of the international application:*	
X the international application as originally filed	
X the description	
pages 1-9 pages NONE	, as originally filed
pages	filed with the domain d
pages NONE filed with the letter of	f
X the claims	
pages 10 NONE	as organish, filo.
as amended (together	with any statement) under Artisle 19
	421.2
pages NONE filed with the letter of	
X the drawings:	
Page NONE	
NONE	as originally filed
pages NONE . filed with the letter of _	filed with the demand
[X] the sequence listing part of the description:	
pages 1-9 pages NONE	as originally filed
pages NONE filed with the letter of	
the language of publication of the international application (under Rule)	48.3(b)).
the language of the translation furnished for the purposes of international prelim or 55.3).	ninary examination (under Rules 55.2 and
With regard to any <b>nucleotide and/or amino acid sequence</b> disclosed in the inte-	arrotional and I also
preliminary examination was carried out on the basis of the sequence listing:	ernational application, the international
X contained in the international application in printed form.	
filed together with the international at the filed together with t	
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furnished subsequently to this Authority in written form.	
furnished subsequently to this Authority in computer readable form.	
The statement that the subsequently furnished written sequence listing does international application as filed has been furnished.	not go beyond the disclosure in the
The statement that the information recorded in computer readable form is identic been furnished.	cal to the writen sequence listing has
X The amendments have resulted in the cancellation of:	,
X the description, pages NONE	
the claims. Nos. NONE	
X the drawings, sheets/Fig NONE	
This report has been drawn as if (some of) the amendments had not been made, so beyond the disclosure as filed, as indicated in the Supplemental Box (Rule 70.2cc	ince they have been considered to go
Replacement sheets which have been furnished to the most supplemental Box (Rule 70.2(c	<u>:)) **</u>
in this report as "originally filed" and are not annexed to this report since they do r and 70.17).	vacaion unaer Article 14 are referred to
and 70.17).  *Any replacement sheet containing such amendments must be referred to under item.	noi contain amenaments (Rules 70.16

## INTERNATIONAL PRELIMINARY EXAMINATION REPORT

International application No

PCT/US00/19007

Novelty (N)	Claims	1-8	YE
	Claims	NONE	NO
Inventive Step (IS)	Claims	1-8	YE
	Claims	NONE	
Industrial Applicability (IA)	Claims	1-8	
radastrai Applicaomy (1.1)	Claims	NONE	YF
NEW CITATIONS			

### INTERNATIONAL PRELIMINARY EXAMINATION REPORT

International application No.

PCT USoo, 19007

VIII.	Certain	observations	on	the	international	application
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The following observations on the clarity of the claims, description, and drawings or on the question whether the claims are fully supported by the description, are made:

Claims 3 and 4 are objected to under PCT Rule 66.2(a)(v) as lacking clarity under PCT Article 6 because the claims are indefinite for the following reason(s). Claims 3 and 4 refer back to a single claim plurals. Appropriate correction is required.

## PATENT COOPERATION TREATY

From the INTERNATIONAL SEARCHING AUTHORITY

To: PAUL J. WHITE NATIONAL RENEWABLE ENERGY LABORATORY 1617 COLE BOULEVARD GOLDEN, CO 80401	PCT
Received	NOTIFICATION OF TRANSMITTAL OF THE INTERNATIONAL SEARCH REPORT
007 <b>2</b> 0 <b>2000</b>	OR THE DECLARATION
Legal Office	(PCT Rule 44.1)
	Date of Mailing (day/month/year) 19 OCT 2000
Applicant's or agent's file reference	(adymorabyear) 10 OCT 2000
PCT/99-45	FOR FURTHER ACTION See paragraphs 1 and 4 below
International application No.	International filing date
PCT/US00/19007	(day/month/year) 13 JULY 2000
MIDWEST RESEARCH INSTITUTE  1. X The applicant is hereby posified that the instance is a second seco	
The applicant is entitled, if he so wishes, to amend the When? The time limit for filing such amendme international search report; however, for m Where? Directly to the International Bureau of WI 34, chemin des Colombette 1211 Geneva 20, Switzerla Facsimile No.: (41-22) 740  For more detailed instructions, see the notes on the Article 17(2)(a) to that effect is transmitted herewith.  3. With regard to the protest against payment of (an) additional services and the second services are such as the second services and the second services are such as the second services and the second services are services as the second second services are services as the second services are services as the second services are services as the second second second services are second	e claims of the international application (see Rule 46):  ints is normally 2 months from the date of transmittal of the more details, see the notes on the accompanying sheet.  PP() es and 1.14.35 he accompanying sheet.  search report will be established and that the declaration under diditional fee(s) under Rule 40.2, the applicant is notified that:
applicant's request to forward the texts of both the	been transmitted to the International Bureau together with the
no decision has been made yet on the protest; the	e applicant will be notified as soon as a decision is made.
4. Further action(s): The applicant is reminded of the follow	ing:
priority claim, must reach the International Bureau as pro completion of the technical preparations for international r	al application will be published by the International Bureau. If notice of withdrawal of the international application, or of the ovided in rules 90 <i>bis</i> 1 and 90 <i>bis</i> 3, respectively, before the publication.
Within 19 months from the priority date, a demand for intern wishes to postpone the entry into the national phase until.	national preliminary examination must be filed if the applicant
	form the prescribed acts for entry into the national phase before

Name and mailing address of the ISA/US Commissioner of Patents and Trademarks Box PCT Washington, D.C 20231

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Authorized officer

MANJUNATH RAO



## **PCT**

### INTERNATIONAL SEARCH REPORT

(PCT Article 18 and Rules 43 and 44)

Applicant's or agent's file reference PCT/99-45	FOR FURTHER ACTION	see Notification of Transmittal of International Search Repo (Form PCT/ISA/220) as well as, where applicable, item 5 below				
International application No. PCT/US00/19007	International filing date 13 JULY 2000	e (day/month/year)	(Earliest) Priority Date (day/month/year) 13 JULY 1999			
Applicant MIDWEST RESEARCH INSTITUTE						
This international search report has been according to Article 18. A copy is being	en prepared by this Internating transmitted to the Intern	ational Searching Au ational Bureau.	thority and is transmitted to the applicant			
This international search report consist  X It is also accompanied by a companied		ument cited in this re	eport.			
1. Basis of the report						
a. With regard to the language, t	he international search was	carried out on the ba	asis of the international application in the			
language in which it was filed the international search was Authority (Rule 23.1(b)).	carried out on the basis	of a translation of the	e international application furnished to this			
	and/or amino acid seque the sequence listing:	nce disclosed in the	international application, the international search			
X contained in the internation	al application in written fo	rm.				
X filed together with the inter	national application in con	nputer readable form				
furnished subsequently to th	is Authority in written for	m.				
furnished subsequently to th	is Authority in computer r	eadable form.				
the statement that the subsc	equently furnished written	sequence listing do	es not go beyond the disclosure in			
the statement that the informa furnished.	tion recorded in computer re	eadable form is identi	cal to the written sequence listing has been			
Certain claims were found		).				
<ul><li>Unity of invention is lacking</li><li>With regard to the title,</li></ul>	ig (See Box II).					
X the text is approved as subm	sitted by the applicant					
the text has been established	• •	as follows:				
5. With regard to the abstract.						
	itted by the applicant.					
the text has been established Box III. The applicant may,	the text is approved as submitted by the applicant.  the text has been established, according to Rule 38.2(b), by this Authority as it appears in Box III. The applicant may, within one month from the date of mailing of this international search report, submit comments to this Authority.					
6. The figure of the <b>drawings</b> to be pu	blished with the abstract is	s Figure No				
as suggested by the applican			None of the figures.			
because the applicant failed						
because this figure better cha	aracterizes the invention.					

### INTERNATION. SEARCH REPORT

ational application No. PCT/US00/19007

A CLA	SCIETO ATION OF CUDITION ATTER					
A. CLASSIFICATION OF SUBJECT MATTER  JPC(7) ::C12N 15/00						
US CL :435/440						
According to International Patent Classification (IPC) or to both national classification and IPC						
	LDS SEARCHED					
	documentation searched (classification system follow-	ed by classification symbols)				
U.S. :	435/440, 435/209, 510 320					
Documenta	ition searched other than minimum documentation to the	he extent that such documents are included	in the fields courshad			
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C. DOC	UMENTS CONSIDERED TO BE RELEVANT					
Category*	Citation of document, with indication, where a	ppropriate, of the relevant passages	Relevant to claim No.			
Y, E	US 6,114,296 A (SCHULEIN ET	AL.) 05 September 2000	1-8			
	(05.09.00), see entire document.	,           •				
• •						
Y	US 5,298,405 A (NEVALAINEN	ET AL.) 29 March 1994	1-8			
	(29.03.94), see entire document.					
Y	US 4,472,504 A (GALLO) 18 Septemb	or 1094 (18 00 04) and anti-	1.0			
1	document.	ger 1984 (18.09.94), see entire	1-8			
i			i			
Y	EP 0,133,035 A2 (SHIN NENRYC	YU KAIHATSU GIJUTSU	1-8			
	KENKYU KUMIAI) 13 February	1985 (13.02.85) see entire				
	document.					
Furth	ner documents are listed in the continuation of Box C	See patent family annex.				
	ecial categories of cited documents					
'A" doc	cument defining the general state of the art which is not considered	"I" later document published after the inte date and not in conflict with the applica principle or theory underlying the inve	tion but cited to understand the			
	be of particular relevance lier document published on or after the international filing date	"X" document of particular relevance, the	i			
'L" doc	cument which may throw doubts on priority claim(s) or which is	considered novel or cannot be consider when the document is taken alone	ed to involve an inventive step			
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'O" doc	cument referring to an oral disclosure lise, exhibition or other means	considered to involve an inventive combined with one or more other such	documents, such combination			
'P" doc the	cument published prior to the international filing date but later than priority date claimed.	being obvious to a person skilled in the "&" document member of the same parent	}			
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Box PCT	ner of Patents and Trademarks	MANJUNATH RAO	(N) UKm			
~	i, D.C. 20231 o. (703) 305-3230	Telephone No. (703) 308-0196				

## (12) INTERNATIONAL APPLICATION PUBLISHED UNDER THE PATENT COOPERATION TREATY (PCT)

## (19) World Intellectual Property Organization International Bureau



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(71) Applicant (for all designated States except US): MID-WEST RESEARCH INSTITUTE [US/US]; 425 Volker Boulevard, Kansas City, MO 64110 (US).

(72) Inventors; and

(75) Inventors/Applicants (for US only): ADNEY, William, S. [US/US]; 13190 West 21st Avenue, Golden, CO 80401 (US). DECKER, Stephen, R. [US/US]; 820 Greenwood Drive, Berthoud, CO 80513 (US). LANTZ McCARTER, Suzanne [US/US]; 3072 West 39th Avenue, Denver, CO 80211 (US). BAKER, John, O. [US/US]; 18790 West 60th Avenue, Golden, CO 80403 (US). NIEVES, Rafael [US/US]; 1794 South Endicott Street, Lakewood, CO

80401 (US). HIMMEL, Michael, E. [US/US]; 9202 West Hialeah Place, Littleton, CO 80123-2148 (US). VINZANT, Todd, B. [US/US]; 16601 W. 15th Avenue, Golden, CO 80401 (US).

(74) Agent: WHITE, Paul, J.; National Renewable Energy Laboratory, 1617 Cole Boulevard, Golden, CO 80401 (US).

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### Published:

With international search report.

For two-letter codes and other abbreviations, refer to the "Guidance Notes on Codes and Abbreviations" appearing at the beginning of each regular issue of the PCT Gazette.

(54) Title: CELLOBIOHYDROLASE REDUCED GLYCOSYLATION VARIANTS: CBHIN45A; CBHIN270A; AND CBHIN384A

(57) Abstract: The invention provides a method for making an active exoglucanase in a eukaryotic heterologous host, the method comprising reducing glycosylation of the exoglucanase, wherein reducing comprises replacing an N-glycosylation site amino acid residue with non-glycosyl accepting amino acid residue. The invention further provides a cellobiohydrolase, comprising the reduced glycosylation variant cellobiose enzymes CBHIN45A; CBHIN270A; or CBHIN384A, or any combination thereof.

WO 01/04284 PCT/US00/19007

# CELLOBIOHYDROLASE REDUCED GLYCOSYLATION VARIANTS: CBHIN45A; CBHIN270A; AND CBHIN384A.

### Technical Field.

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This invention relates to exoglucanases. More specifically, it relates to *Trichoderma* reesei cellobiohydrolase I reduced glycosylation variants which enable expression of the active enzyme in a heterologous host.

### Background Art.

The surface chemistry of acid pretreated-biomass, used in ethanol production, is different from that found in plant tissues, naturally digested by fungal cellulase enzymes, in two important ways: (1) pretreatment heats the substrate past the phase-transition temperature of lignin; and (2) pretreated biomass contains less acetylated hemicellulose. Thus, it is believed, that the cellulose fibers of pretreated-biomass are coated with displaced and modified lignin. This alteration results in a non-specific binding of the protein with the biomass, which impedes enzymatic activity. Moreover, where the pretreated biomass is a hardwood-pulp it contains a weak net-negatively charged surface, which is not observed in native wood. Therefore, for the efficient production of ethanol from pretreated biomass it is desirable to enhance the catalytic activity of glycosyl hydrolases on acid hydrolyzed hardwoods.

Trichoderma reesei CBH I is a mesophilic cellulase enzyme, and comprises a major catalyst in the overall hydrolysis of cellulose. An artificial ternary cellulase system consisting of a 90:10:2 mixture of *T. reesei* CBH I, *A. cellulolyticus* EI, and *A. niger* β-D-glucosidase is capable of releasing as much reducing sugar from pretreated yellow poplar as the native *T. reesei* system after 120 h. This result is encouraging for the ultimate success of engineered cellulase systems, because this artificial enzyme system was tested at 50°C, a temperature far below that considered optimal for EI, in order to spare the more heat labile enzymes CBH I and β-D-glucosidase. In order to increase the efficiency of such artificial enzyme systems it is desirable to engineer new *T. reesei* CBH I variant enzymes capable of active expression in a heterologous host. The heterologous host *Aspergillus awamori*, could provide an excellent capacity for synthesis and secretion of *T. reesei* CBH I because of its ability to correctly fold and post-translationally modify proteins of eukaryotic origin. Moreover, *A. awamori* is believed to be an excellent test-bed for *Trichoderma* coding sequences and resolves some of the problems associated with direct site directed mutagenesis in *Trichoderma*.

In consideration of the foregoing, it is therefore desirable to provide variant cellulase

enzymes having enzymatic activity when expressed in an heterologous host.

### Disclosure of Invention.

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It is a general object of the present invention to provide variant cellulase enzymes having enzymatic activity when expressed in a heterologous host, such as a filamentous fungi or yeast.

Another object of the invention is to provide a variant exoglucanase characterized by a reduction in glycosylation when expressed in a heterologous host.

Another object of the invention is to provide an active cellobiohydrolase enzyme capable of expression in heterologous fungi or yeast.

It is yet another object of the invention to provide a method for reducing the glycosylation of a cellobiohydrolase enzyme for expression in a heterologous host.

The foregoing specific objects and advantages of the invention are illustrative of those which can be achieved by the present invention and are not intended to be exhaustive or limiting of the possible advantages which can be realized. Thus, those and other objects and advantages of the invention will be apparent from the description herein or can be learned from practicing the invention, both as embodied herein or as modified in view of any variations which may be apparent to those skilled in the art.

Briefly, the invention provides a method for making an active exoglucanase in a heterologous host, the method comprising reducing glycosylation of the exoglucanase, reducing glycosylation further comprising replacing an N-glycosylation site amino acid residue with a non-glycosyl accepting amino acid residue. The invention further provides a cellobiohydrolase, comprising the reduced glycosylation variant cellobiose enzymes CBHIN45A; CBHIN270A; or CBHIN384A, or any combination thereof.

### 25 Best Mode for Carrying out the Invention.

Unless specifically defined otherwise, all technical or scientific terms used herein have the same meaning as commonly understood by one of ordinary skill in the art to which this invention belongs. Although any methods and materials similar or equivalent to those described herein can be used in the practice or testing of the present invention, the preferred methods and materials are now described.

A method for reducing the glycosylation of an expressed *Trichoderma reesei* CBHI protein by site-directed mutagenesis ("SDM") is disclosed. The method includes replacing an N-glycosylation site amino acid residue, such as asparagines 45, 270, and/or 384 of SEQ. ID NO: 4

(referenced herein as CBHIN45A, CBHIN270A and CBHIN384A, respectively), with a non-glycosyl accepting amino acid residue, such as is alanine. Various mutagenesis kits for SDM are available to those skilled in the art and the methods for SDM are well known. The description below discloses a procedure for making and using CBHI variants: CBHIN45A; CBHIN270A; and CBHIN384A. The examples below demonstrate the expression of active CBH I in the heterologous fungus *Aspergillus awamori*.

### Industrial Applicability

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Site-Directed Mutagenesis of Trichoderma reesei CBH I for Reduced Glycosylation.

Aspergillus awamori was transformed with various versions of the cbhl gene from Trichoderma reesei. The cbhl genes included both cDNA and genomic (intron containing) versions. These were altered by site-directed mutagenesis for the specific purpose of reducing the glycosylation of the expressed CBH I protein through replacement of the N-glycosylation site amino acid residues (asparagine) with non-glycosyl accepting amino acid residues (alanine). The gene was propagated in an E. coli vector plasmid (pPFE2) under the control of the Aspergillus awamori glucoamylase promoter and signal sequence, and trpC terminator, and carrying resistance to ampicillin (E. coli selection) and Zeocin (Bleomycin) Aspergillus selection. One altered rCBH I variant, CBHIN270A, SEQ. ID. NO: 2, was isolated from cultures and determined to be consistent with native CBH I, SEQ. ID. NO: 4, with respect to kinetics on pNPL and was only slightly higher in molecular weight. Thus, construction of the triple reduced glycosylation mutant CBH 1, CBHIN270A (SEQ. ID. NO: 2) / CBHIN45A (SEQ. ID. NO: 1) / CBHIN384A (SEQ. ID. NO: 3), may provide a viable means of producing active CBH I in heterologous fungal or yeast which do not require the cellobiose/lactose induction cascade, known in Trichoderma. It is believed that reduced glycosylation CBH I mutants would also serve effectively in yeast-based high throughput screens, which are normally rendered unusable for fungal enzymes because of hyperglycosylation.

## Example 1. Production of Active Recombinant CBH I (rCBH I) in Aspergillus awamori Construction of Modified CBH I Coding Sequence.

The coding sequence for *T. reesei* CBH I (SEQ. ID. NO: 4) was successfully inserted and expressed in *Aspergillus awamori* using the fungal expression vector pPFE2 (and pPFE1). Vectors pPFE1 and pPFE2 are *E. coli-Aspergillus* shuttle vectors, and contain elements required for maintenance in both hosts. They encode ampicillin resistance for selection in *E. coli* and

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Zeocin resistance for selection in Aspergillus. The foregoing provided for the site-directed mutagenesis in E. coli, followed by expression of the new mutant proteins in A. awamori. The CBH I gene is under the control of the A. awamori glucoamylase promoter and includes the glucoamylase secretion signal peptide. In order to have the signal peptide properly cleaved during secretion, the construction of this plasmid required the addition, by PCR, of a Notl site and Xbal site on the coding sequence of CBH 1. The Notl site addition resulted in a change of the most N-terminal amino acid on the protein from glutamine to glycine. This glycine was subsequently changed back to the native glutamine in the pPFE2/CBHI construct, using site-directed mutagenesis PCR. This new construct was used to transform A. awamori and to express rCBH I, as confirmed by western blot analysis of culture supernatant. The rCBH I expressed in A. awamori tends to be over glycosylated as evidenced by the higher molecular weight observed on western blot analysis. Over-glycosylation of CBH I by A. awamori was confirmed by digestion of the recombinant protein with endoglycosidases. Following endoglycosidase H and F digestion, the higher molecular weight form of the protein collapses to a molecular weight similar to native CBH 1.

The vector pPFE2/CBHI requires a relatively long PCR reaction (8.2 kb) to make site-specific changes using the Stratagene Quik Change protocol. The PCR reaction was optimized as follows using a GeneAmp PCR System 2400, Perkin Elmer Corporation. The reaction mixture contained 50 ng of template DNA, 125 ng each of the sense and antisense mutagenic primers, 5 µl of Stratagene 10x cloned Pfu buffer, 200µM of each: dNTP, 5 mM MgCl<sub>2</sub> (total final concentration Of MgCl<sub>2</sub> is 7 mM); and 2.5 U Pfu Turbo DNA polymerase. The PCR reaction was carried out for 30 cycles, each consisting of one minute denaturation at 96°C, 1 minute annealing at 69°C, and 20-minute extension at 75°C. There is an initial denaturation for 2 minutes at 96°C and a final extension for 10 minutes at 75°C, followed by a hold at 4°C. Agarose gel electrophoresis, ethidium bromide staining, and visualization under UV transillumination were used to confirm the presence of a PCR product.

PCR products were digested with restriction enzyme Dpnl, to degrade un-mutagenized parental DNA, and transformed into *E. coli* (Stratagene Epicurian Coli Supercompetent XL-1 Cells). Amp<sup>R</sup> colonies were picked from LB-Amp<sup>100</sup> plates and mutations were confirmed by DNA sequencing. Depending on scale, plasmid DNA was purified using the Qiagen QiaPrep Spin Miniprep Kit or the Promega Wizard Plus MaxiPrep DNA Purification System.

Transformation of Aspergillus awamori with Trichoderma reesei CBH I coding sequence.

Aspergillus awamori spore stocks were stored at -70°C in 20% glycerol 10% lactose. After thawing, 200 µL of spores were inoculated into 50 mL CM broth in each of eight baffled 250 mL Erlenmeyer flask. The cultures were grown at 28°C, 225 rpm for 48 h. The mycelia were removed by filtration with sterile Miracloth, Calbiochem, San Diego, CA, and washed thoroughly with sterile KCM. Approximately 10 g of washed mycelia were transferred to 50 mL KCM + 250 mg Novozym234 in a 250 mL baffled Erlenmeyer flask. The digestion mixture was incubated at 30°C, 80 rpm for 16-18 h. Spheroplasts were filtered through Miracloth into 50 mL conical centrifuge tubes, pelleted at 2000xg for 15 min and re-suspended in 0.7M KCl by gentle tituration with a 25 mL pipette. This procedure was repeated once. After a third pelleting, the spheroplasts were resuspended in 10 mL KC, pelleted and resuspended in 0.5 mL KC using a wide-bore pipet tip. The washed spheroplasts were transformed by adding 12.5  $\mu$ L PCM and 5 uL DNA (≈0.5 ug/uL) to 50 uL of spheroplast in sterile 1.5 mL Eppendorf tubes. After incubation on ice for 45 minutes, 0.5 mL of room temperature PCM was added to the transformation mixture and was mixed by tituration with a wide bore pipet tip. The mixture was incubated at room temperature for 45 minutes. One milliliter of KC was added and mixed. The mixture was allocated between four tubes of molten CM top agar at 55°C, which were each poured over a 15 mL CM170 plate. The plates were incubated at 28°C for 2-3 days. Subsurface colonies were partially picked with a sterile wide bore pipet tip, exposing the remaining part of the colony to air and promoting rapid sporulation. After sporulation, spores were streaked onto several successive CM100 plates. After a monoculture was established, heavily sporulated plates were flooded with sterile spore suspension medium (20% glycerol, 10% lactose), the spores were suspended and aliquots were frozen at -70°C. Protein production was confirmed and followed by western blot using anti-CBH I monoclonal antibodies and the Novex Western Breeze anti-mouse chromogenic detection kit (Novex, San Diego, CA). Extracting genomic DNA using the YeaStar Genomic DNA Kit (Zymo Research, Orange, CA) and carrying out PCR with pfu-turbo DNA polymerase (Stratagene, La Jolla, CA) and cbhl primers confirmed insertion of the gene

### Production of Recombinant Enzyme.

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For enzyme production, spores were inoculated into 50 mL CM maltose medium, pH 5.0, and grown at 32°C, 225 rpm in 250 mL baffled flasks. The cultures were transferred to 1.0 L of CM maltose in 2,800 mL Fernbach flasks and grown under similar conditions. For large-scale enzyme production (>1 mg), these cultures were transferred to 10-L CM maltose in a New Brunswick BioFlo3000 chemostat (10-L working volume) maintained at: 20% DO; pH 4.5; 25°C;

and 300 rpm. The culture was harvested by filtration through Miracloth after 2-3 days of growth. For the 10-L fermentation broth, the filtrate was concentrated and dia-filtered into 20mM sodium acetate pH 5.0 by tangential flow ultrafiltration with an Amicon DC30 concentrator equipped with a single 10,000 MWCO hollow fiber cartridge (l.1mm I.D., 2.4 m² surface area). The retentate from the 10-L concentration or the filtrate from smaller cultures was clarified in an Amicon DC-2 concentrator by tangential flow filtration with two 0.1 µm hollow fiber cartridges (1.1 mm I.D., 0.03 m² surface area, Millipore, Bedford, MA). The permeate was further concentrated with an Amicon CH-2 concentrator equipped with three 10,000 MWCO hollow fiber cartridges (1.1 mm I.D., 0.03 m² surface area). The final concentrate was sterile filtered through a 0.45 µm filter and stored at 4°C until used.

The recombinant CBH I protein, SEQ. ID NO.: 4, was purified by passing the concentrated culture broth over two or three CBinD900 cartridge columns (Novagen, Madison, WI) connected in series using a Pharmacia FPLC loading at 1.0 mL/min. (Amersham Pharmacia Biotech, Inc., Piscataway, NJ). The cartridges were equilibrated in 20 mM Bis-Tris pH 6.5 prior to loading and washed with the same buffer after loading. The bound rCBH I was eluted with 100% ethylene glycol (3 mL/column) by hand, using a syringe. The eluted rCBH I was concentrated in an Amicon 10 mL stirred cell using a 25 mm PM10 membrane to <2.0 mL and loaded onto a Pharmacia SuperDex200 16/60 size-exclusion column. The mobile phase was: 20 mM sodium acetate; 100 mM sodium chloride; and 0.02% sodium azide, pH 5.0 running at 1.0 mL/min. The eluted protein was concentrated by stirred cell and stored at 4°C. Concentration was determined by A<sub>280</sub> using the extinction coefficient and molecular weight calculated for individual proteins by the ProtParam tool on the ExPASy website (http://expasy.ch/tools/protparam.html). Below are the formulations for the various media described herein:

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### Clutterbuck's Salts (20X)

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Na <sub>2</sub> NO <sub>3</sub>	120.0 g/L
KCl	10.4 g/L
$MgSO_4*7H_20$	10.4 g/L
KH <sub>2</sub> PO <sub>4</sub>	30.4 g/L
CM- Yeast Extract-	5g/L
Tryptone-	5g/L
Glucose-	10g/L

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Clutterbuck's Salts-50mL

Add above to 900mL dH<sub>2</sub>0, pH to 7.5, bring to 1000mL

CM Agar- CM+ 20g/L Agar

CMK CM Agar+ 0.7M KCl

CM100- CM + 100 g/mL Zeocin (Invitrogen, Carlsbad, CA)

CM 1070- CM+ 170 g/mL Zeocin

KCl- 0.7M KCl

KC- 0.7M KCl + 50mM CaCI2

KCM- 0.7M KCl + 10mM MOPS, pH 5.8

PCM 40% PEG 8000, 50mM CaCl<sub>2</sub>, 10mM MOPS pH 5.8

### Example 2. Production of Reduced Glycosylation rCBH 1: Sites N270A; N45A; and N384A.

rCBHI/pPFE2 has been optimized using site-directed mutagenesis to achieve expression of native molecular weight CBH I in A. awamori. The QuickChange SDM kit (Stratagene, San Diego, Ca) was used to make point mutations, switch amino acids, and delete or insert amino acids in the native CBH1 gene sequence. The Quick Change SDM technique was performed using thermotolerant Pfu DNA polymerase, which replicates both plasmid strands with high fidelity and without displacing the mutant oligonucleotide primers. The procedure used the polymerase chain reaction (PCR) to modify the cloned CBH1 DNA. The basic procedure used a supercoiled double stranded DNA (dsDNA) vector, with an insert of interest, and two synthetic oligonucleotide primers containing a desired mutation. The oligonucleotide primers, each complimentary to opposite strands of the vector, extend during temperature cycling by means of the polymerase. On incorporation of the primers, a mutated plasmid containing staggered nicks was generated. Following temperature cycling, the product was treated with a Dpn1 restriction enzyme. Dpn1 is specific for methylated and hemi-methylated DNA and thus digests the unmutated parental DNA template, selecting for the mutation-containing, newly-synthesized DNA. The nicked vector DNA, containing the desired mutations, was then transformed into E. coli. The small amount of template DNA required to perform this reaction, and the high fidelity of the Pfu DNA polymerase contribute to the high mutation efficiency and minimizes the potential for the introduction of random mutations. Three glycosylation-site amino acids on the protein surface were targeted for substitution of an alanine (A) residue in place of asparagine (N). Single site substitutions were successfully completed in the CBH I coding sequence at sites N45, N270, and N384, of SEQ. ID NO.: 4 by site-directed mutagenesis, and confirmed by DNA sequencing.

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Table 1.

CONSTRUCT	HOST	MW	K <sub>M</sub>	$V_{MAX}$
T. reesei	none	57.8 kDa	1.94	0.746
rCBH I wt cDNA#	A. awamori	63.3 kDa	2.14	0.668
rCBH I wt genomic	A. awamori	63.3 kDa		
rCBH I N270A	A. awamori	61.7 kDa	2.25	0.489

As shown in **Table 1**, Western blot analysis of the supernatant, obtained from a single glycosylation-site mutant CBHIN270A (SEQ. ID NO.: 2) culture expressed in *A. awamori*, demonstrated that a decrease, to a lower molecular weight (61.7 kDa), in the amount of protein had occurred, as compared to the that in the wild type cDNA (63.3 kDa), and the wild type genomic DNA (63.3 kDa). These results demonstrate a reduction in the level of glycosylation in the reduced glycosylation mutant CBHIN270A, via expression in *A. awamori*. It is also shown, in the Table, that the CBHIN270A enzyme nearly retained its native enzymatic activity when assayed using the *p*NPL substrate. While not shown in the Table, variants CBHIN45A (SEQ. ID NO.: 1), and CBHI384A (SEQ. ID NO.: 3) have also demonstrated a reduction in amount of glycosylation and native activity when expressed from the heterologous host *A. awamori*.

## 20 Example 3. Production of Reduced Glycosylation rCBH 1: Double and Triple Mutants.

Double and triple combinations of this substitution have also been completed in the CBH I coding sequence (SEQ. ID NO.: 4) at sites N45, N270, and N384 by site-directed mutagenesis and confirmed by DNA sequencing. These double and triple-site constructs will also yield rCBH I enzymes with reduced glycosylation and, presumably, native activity.

### Mutagenic Primers Used in Site-directed Mutagenesis PCR

Not1, Xbal insertion for vector construction

Mutagenic primers

C-terminal strand (Xbal): AGAGAGTCTAGACACGGAGCTTACAGGC

N-terminal strand (Notl):

AAAGAAGCGCGGCCGCCTGCACTCTCCAATCGG

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Repair of Notl site to native sequence

Mutagenic primers

sense strand:GGCAAATGTGATTTCCAAGCGCCAGTCGGCCTGCACTCTCC

antisense strand:GGAGAGTGCAGGCCGACTGGCGCTTGGAAATCACATTTGCC

N45A glycosylation site mutation

Mutagenic primers

sense strand- GGACTCACGCTACGGCCAGCAGCACGAACTGC antisense strand: GCAGTTCGTGCTGCTGCCGTAGCGTGAGTCC

N270A glycosylation site mutation

Mutagenic primers

sense strand: CCCATACCGCCTGGGCGCCACCAGCTTCTACGGCCC

antisense strand: GGGCCGTAGAAGCTGGTGGCGCCCAGGCGGTATGGG

N384A glycosylation site mutation

Mutagenic primers

sense strand: GGACTCCACCTACCCGACAGCCGAGACCTCCTCCACACCCG

20 antisense strand:

CGGGTGTGGAGGAGGTCTCGGCTGTCGGGTAGGTGGAGTCC

The foregoing description is considered as illustrative only of the principles of the invention. Furthermore, since numerous modifications and changes will readily occur to those skilled in the art, it is not desired to limit the invention to the exact construction and process shown as described above. Accordingly, all suitable modifications and equivalents may be resorted to falling within the scope of the invention as defined by the claims which follow.

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10 Claims

- 1. A method for making an active exoglucanase in a eukaryotic heterologous host, the method comprising reducing glycosylation of the exoglucanase, wherein reducing comprises replacing an N-glycosylation site amino acid residue with non-glycosyl accepting amino acid residue.
- 2. The method of claim 1, wherein the N-glycosylation site amino acid residues include asparagine 45, 270, or 384 of SEQ ID NO: 4 and the non-glycosyl accepting amino acid residue includes alanine.
  - 3. The method of claims 1 wherein replacing comprises site-directed-mutagenesis.
  - 4. The methods of claims 1 wherein the exoglucanase comprises a cellobiohydrolase.
  - 5. An exoglucanase, comprising SEQ. ID. NO: 1.
  - 6. An exoglucanase, comprising SEQ. ID. NO: 2.
  - 7. An exoglucanase, comprising SEQ. ID. NO: 3.
  - 8. An exoglucanase, comprising a combination of claims 5,6, or 7.

### **SEQUENCE LISTING**

<110> Adney, William S. Decker, Stephen R. Lantz-McCarter, Suzanne Baker, John O. Vinzant, Todd B. Nieves, Rafael A Himmel, Michael E. <120> CELLOBIOHYDROLASE REDUCED GLYCOSYLATION VARIANTS: CBHIN45A; CBHIN270A; AND CBHIN384A <130> HIMMEL NREL IR# 99-45 <140> <141> <160>4 <170> PatentIn Ver. 2.0 <210> 1 <211> 496 <212> PRT <213> Trichoderma reesei <400> 1 Gln Ser Ala Cys Thr Leu Gln Ser Glu Thr His Pro Pro Leu Thr Trp 10 5 Gln Lys Cys Ser Ser Gly Gly Thr Cys Thr Gln Gln Thr Gly Ser Val 25 Val Ile Asp Ala Asn Trp Arg Trp Thr His Ala Thr Ala Ser Ser Thr 35 40 Asn Cys Tyr Asp Gly Asn Thr Trp Ser Ser Thr Leu Cys Pro Asp Asn Glu Thr Cys Ala Lys Asn Cys Cys Leu Asp Gly Ala Ala Tyr Ala Ser 75 Thr Tyr Gly Val Thr Thr Ser Gly Asn Ser Leu Ser Ile Gly Phe Val Thr Gln Ser Ala Gln Lys Asn Val Gly Ala Arg Leu Tyr Leu Met Ala 110 100

Ser Asp Thr Thr Tyr Gln Glu Phe Thr Leu Leu Gly Asn Glu Phe Ser

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			2		
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Tyr Phe Va	l Ser Met Asp	Ala Asp Gly	Gly Val Ser 155		<b>hr</b> 50
Asn Thr Ala	a Gly Ala Lys 7	Tyr Gly Thr C	Gly Tyr Cys A	Asp Ser Gln C	
	103	i	70	175	
Pro Arg As	p Leu Lys Phe 180	Ile Asn Gly ( 185	Hn Ala Asn	Val Glu Gly Ti 190	пр
Glu Pro Ser 195	Ser Asn Asn A	Ala Asn Thr ( 200		lly His Gly Ser 05	
Cys Cys Ser 210	Glu Met Asp	lle Trp Glu A 215	Ja Asn Ser I 220	le Ser Glu Ala	
Leu Thr Pro 225	His Pro Cys T 230	hr Thr Val G	Hy Gln Glu I 235	le Cys Glu Gly 240	
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Asp Pro Asp	Gly Cys Asp 7 260	Frp Asn Pro ' 265	Гуг Arg Leu	Gly Asn Thr S	Ser
Phe Tyr Gly 275	Pro Gly Ser Se	er Phe Thr Le 280		hr Lys Lys Le 85	u
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Val Gln Asn 305	Gly Val Thr Pl 310	ne Gln Gln Pr	o Asn Ala C	Hu Leu Gly Se 320	
Гуг Ser Gly A	Asn Glu Leu A 325	sn Asp Asp T 3	Tyr Cys Thr .	Ala Glu Glu A 335	la
Giu Phe Gly	Gly Ser Ser Ph 340	e Ser Asp Ly 345	s Gly Gly Le	eu Thr Gln Phe 350	;
Lys Lys Ala 7 355	Thr Ser Gly Gl	y Met Val Le 360		er Leu Trp As 55	p
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Ser Gly Val Pro Ala Gln Val Glu Ser Gln Ser Pro Asn Ala Lys Val 405 410 415

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Ser Gly Gly Asn Pro Pro Gly Gly Asn Arg Gly Thr Thr Thr Arg
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440
445

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Ser Gly Thr Thr Cys Gln Val Leu Asn Pro Tyr Tyr Ser Gln Cys Leu 485 490 495

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Asn Cys Tyr Asp Gly Asn Thr Trp Ser Ser Thr Leu Cys Pro Asp Asn 50 55 60

Glu Thr Cys Ala Lys Asn Cys Cys Leu Asp Gly Ala Ala Tyr Ala Ser 65 70 75 80

Thr Tyr Gly Val Thr Thr Ser Gly Asn Ser Leu Ser Ile Gly Phe Val 85 90 95

Thr Gln Ser Ala Gln Lys Asn Val Gly Ala Arg Leu Tyr Leu Met Ala 00 105 110

Ser Asp Thr Thr Tyr Gln Glu Phe Thr Leu Leu Gly Asn Glu Phe Ser 115 120 125

Phe Asp Val Asp Val Ser Gln Leu Pro Cys Gly Leu Asn Gly Ala Leu

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Asn Thr Ala	Gly Ala Lys T 165	yr Gly Thr	Gly Tyr Cy 170	s Asp Ser	Gln Cys 175
Pro Arg Asp	Leu Lys Phe 1	Ile Asn Gly 185	Gln Ala As	n Val Glu 190	Gly Trp
Glu Pro Ser 195	Ser Asn Asn A	Ala Asn Thr 200	Gly Ile Gly	Gly His C 205	Hy Ser
Cys Cys Ser 210	Glu Met Asp	Ile Trp Glu 215	Ala Asn Se	er Ile Ser G 20	ilu Ala
Leu Thr Pro 225	His Pro Cys T 230	Thr Thr Val	Gly Gln Gl 235	u Ile Cys (	Glu Gly 240
Asp Gly Cys	s Gly Gly Thr 7 245	Гуг Ser Asp	Asn Arg T 250	yr Gly Gly	Thr Cys 255
Asp Pro Asp	Gly Cys Asp 260	Trp Asn Pr 26		Leu Gly Al 27	
Phe Tyr Gly 275	Pro Gly Ser S	er Phe Thr 280	Leu Asp T	hr Thr Lys 285	Lys Leu
Thr Val Val 290	Thr Gln Phe C	Glu Thr Ser 95	Gly Ala Ile 300		Гуг Туг
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Tyr Ser Gly	Asn Glu Leu A	Asn Asp As	p Tyr Cys 7 330	Thr Ala Gl	u Glu Ala 335
Glu Phe Gly	Gly Ser Ser P 340	he Ser Asp 345	Lys Gly Gl	ly Leu Thr 350	Gin Phe
Lys Lys Ala	Thr Ser Gly C	Gly Met Val 360	Leu Val M	let Ser Leu 365	Trp Ası

Asp Tyr Tyr Ala Asn Met Leu Trp Leu Asp Ser Thr Tyr Pro Thr Asn 370 375 380

Glu Thr Ser Ser Thr Pro Gly Ala Val Arg Gly Ser Cys Ser Thr Ser 385 390 395 400

Ser Gly Val Pro Ala Gln Val Glu Ser Gln Ser Pro Asn Ala Lys Val

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Thr Phe Ser Asn Ile Lys Phe Gly Pro Ile Gly Ser Thr Gly Asn Pro 420 425 430

Ser Gly Gly Asn Pro Pro Gly Gly Asn Arg Gly Thr Thr Thr Thr Arg 435 440 445

Arg Pro Ala Thr Thr Gly Ser Ser Pro Gly Pro Thr Gln Ser His 450 455 460

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Asn Cys Tyr Asp Gly Asn Thr Trp Ser Ser Thr Leu Cys Pro Asp Asn 50 55 60

Glu Thr Cys Ala Lys Asn Cys Cys Leu Asp Gly Ala Ala Tyr Ala Ser 70 75 80

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Asn Thr Ala Gly Ala Lys Tyr Gly Thr Gly Tyr Cys Asp Ser Gln Cys 165 170 175

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180 185 190

Glu Pro Ser Ser Asn Asn Ala Asn Thr Gly Ile Gly Gly His Gly Ser 195 200 205

Cys Cys Ser Glu Met Asp Ile Trp Glu Ala Asn Ser Ile Ser Glu Ala 210 215 220

Leu Thr Pro His Pro Cys Thr Thr Val Gly Gln Glu Ile Cys Glu Gly 225 230 235 240

Asp Gly Cys Gly Gly Thr Tyr Ser Asp Asn Arg Tyr Gly Gly Thr Cys 245 250 255

Asp Pro Asp Gly Cys Asp Trp Asn Pro Tyr Arg Leu Gly Asn Thr Ser 260 265 270

Phe Tyr Gly Pro Gly Ser Ser Phe Thr Leu Asp Thr Thr Lys Lys Leu 275 280 285

Thr Val Val Thr Gln Phe Glu Thr Ser Gly Ala Ile Asn Arg Tyr Tyr 290 295 300

Val Gln Asn Gly Val Thr Phe Gln Gln Pro Asn Ala Glu Leu Gly Ser 305 310 315 320

Tyr Ser Gly Asn Glu Leu Asn Asp Asp Tyr Cys Thr Ala Glu Glu Ala 325 330 335

Glu Phe Gly Gly Ser Ser Phe Ser Asp Lys Gly Gly Leu Thr Gln Phe 340 345 350

Lys Lys Ala Thr Ser Gly Gly Met Val Leu Val Met Ser Leu Trp Asp 355 360 365

Asp Tyr Tyr Ala Asn Met Leu Trp Leu Asp Ser Thr Tyr Pro Thr Ala 370 375 380

Glu Thr Ser Ser Thr Pro Gly Ala Val Arg Gly Ser Cys Ser Thr Ser 385 390 395 400

Ser Gly Val Pro Ala Gln Val Glu Ser Gln Ser Pro Asn Ala Lys Val 405 410 415 Thr Phe Ser Asn Ile Lys Phe Gly Pro Ile Gly Ser Thr Gly Asn Pro 420 425 430

Ser Gly Gly Asn Pro Pro Gly Gly Asn Arg Gly Thr Thr Thr Thr Arg 435 440 445

Arg Pro Ala Thr Thr Gly Ser Ser Pro Gly Pro Thr Gln Ser His 450 455 460

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Val Ile Asp Ala Asn Trp Arg Trp Thr His Ala Thr Asn Ser Ser Thr 35 40 45

Asn Cys Tyr Asp Gly Asn Thr Trp Ser Ser Thr Leu Cys Pro Asp Asn 50 55 60

Glu Thr Cys Ala Lys Asn Cys Cys Leu Asp Gly Ala Ala Tyr Ala Ser 70 75 80

Thr Tyr Gly Val Thr Thr Ser Gly Asn Ser Leu Ser Ile Gly Phe Val 85 90 95

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Phe Asp Val Asp Val Ser Gln Leu Pro Cys Gly Leu Asn Gly Ala Leu 130 135 140

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Glu Pro Ser Ser Asn Asn Ala Asn Thr Gly Ile Gly Gly His Gly Ser 195 200 205

Cys Cys Ser Glu Met Asp Ile Trp Glu Ala Asn Ser Ile Ser Glu Ala 210 215 220

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Tyr Ser Gly Asn Glu Leu Asn Asp Asp Tyr Cys Thr Ala Glu Glu Ala 325 330 335

Glu Phe Gly Gly Ser Ser Phe Ser Asp Lys Gly Gly Leu Thr Gln Phe 340 345 350

Lys Lys Ala Thr Ser Gly Gly Met Val Leu Val Met Ser Leu Trp Asp 355 360 365

Asp Tyr Tyr Ala Asn Met Leu Trp Leu Asp Ser Thr Tyr Pro Thr Asn 370 375 380

Glu Thr Ser Ser Thr Pro Gly Ala Val Arg Gly Ser Cys Ser Thr Ser 385 390 395 400

Ser Gly Val Pro Ala Gln Val Glu Ser Gln Ser Pro Asn Ala Lys Val 405 410 415

Thr Phe Ser Asn Ile Lys Phe Gly Pro Ile Gly Ser Thr Gly Asn Pro 420 425 430

Ser Gly Gly Asn Pro Pro Gly Gly Asn Arg Gly Thr Thr Thr Thr Arg 435 440 445

Arg Pro Ala Thr Thr Gly Ser Ser Pro Gly Pro Thr Gln Ser His 450 455 460

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### INTERNATIONAL SEARCH REPORT

International application No. PCT/US00/19007

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A. CLA IPC(7)	SSIFICATION OF SUBJECT MATTER :C12N 15/00					
US CL :435/440  According to International Patent Classification (IPC) or to both national classification and IPC						
B. FIELDS SEARCHED						
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U.S. :	435/440, 435/209, 510/320					
Documental	tion searched other than minimum documentation to the	ne extent that such documents are included	in the fields searched			
	lata base consulted during the international search (n LUS BIOSIS MEDLINE BIOTECHABS	ame of data base and, where practicable,	search terms used)			
C. DOC	UMENTS CONSIDERED TO BE RELEVANT					
Category*	Citation of document, with indication, where a	ppropriate, of the relevant passages	Relevant to claim No.			
Y, E	US 6,114,296 A (SCHULEIN ET (05.09.00), see entire document.	AL.) 05 September 2000	1-8			
Y	US 5,298,405 A (NEVALAINEN ET AL.) 29 March 1994 1-8 (29.03.94), see entire document.					
Y	US 4,472,504 A (GALLO) 18 September 1984 (18.09.94), see entire document.					
Y	EP 0,133,035 A2 (SHIN NENRYO KENKYU KUMIAI) 13 February document.		1-8			
Furth	er documents are listed in the continuation of Box C	. See patent family annex.				
"A" doc	cial categories of cited documents:  ument defining the general state of the art which is not considered be of particular relevance	"T" later document published after the inte- date and not in conflict with the applica principle or theory underlying the inve	tion but cited to understand the			
"L" doc	ier document published on or after the international filing date ument which may throw doubts on priority claim(s) or which is d to establish the publication date of another citation or other	"X" document of particular relevance; the considered novel or cannot be consider when the document is taken alone	claimed invention cannot be ed to involve an inventive step			
-	special reason (as specified)  "Y"  document of particular relevance; the claimed invention cannot be considered to involve an inventive step when the document is					
	ument published prior to the international filing date but later than priority date claimed	being obvious to a person skilled in the "&" document member of the same patent				
	MBER 2000	Date of mailing of the international sear 19 OCT 2000	rch report			
Commission Box PCT	nailing address of the ISA/US her of Patents and Trademarks D.C. 20231	Authorized officer  MANJUNATH RAD	Meykon			
-	o. (703) 305-3230	Telephone No. (703) 308-0196				